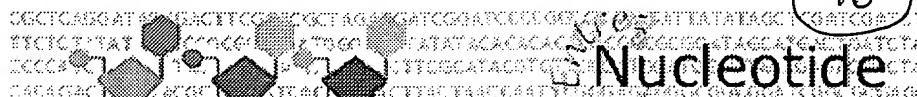


Score = 71.9 bits (36), Expect = 2e-10  
Identities = 63/72 (87%)  
Strand = Plus / Plus

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Sbjct: 132  tcatcaaacaat 143
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SEQ ID NO: 16



Boo

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## Details

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Protein, PubMed, Taxonomy, LinkOut

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BASE COUNT 512 a 334 c 318 g 458 t  
ORIGIN

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Revised: July 5, 2002.

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Jul 31 2002 10:35:46